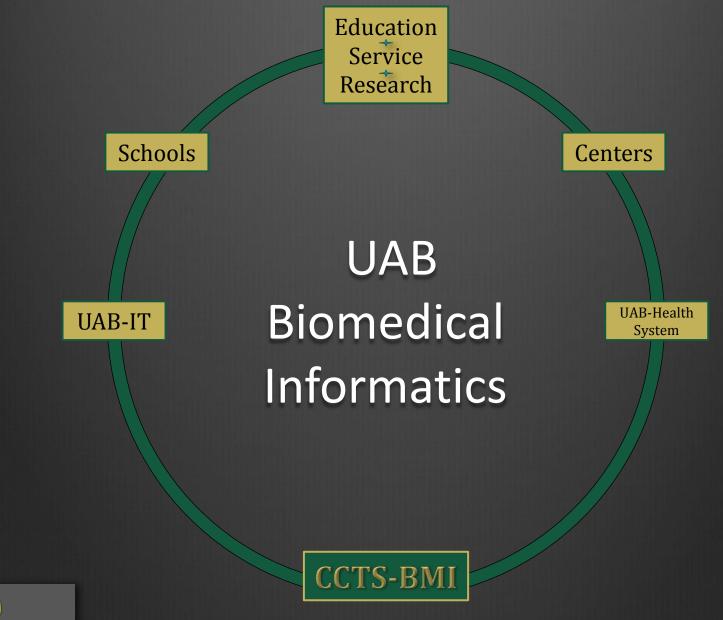
## **Biomedical Informatics**

A Partnership for UAB





# CCTS Biomedical Informatics Component (BMI)

#### > Leadership

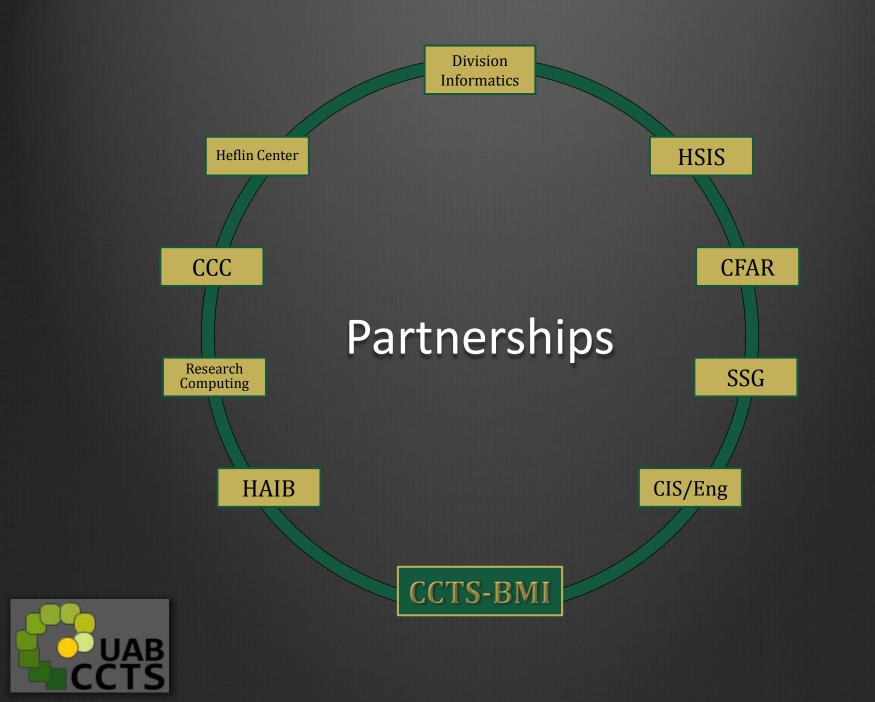
- Elliot Lefkowitz, Director
- Eta Berner, Co-Director
- Jonas Almeida, Co-Director
- Matt Wyatt, Assistant Director



#### ➤ Staff

- Bioinformaticians: David Crossman, Curtis Hendrickson, Ranjit Kumar, John Osborne
- System Analysts: Keith Bobitt, Don Dempsey, Nivi Totha

### http://bioinformatics.uab.edu



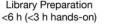
# Heflin Center for Genomic Sciences Core Laboratories

Molly Bray, PhD Michael Crowley, PhD David Crossman, PhD

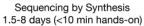
## Next Generation Sequencing

- The Heflin Genomics Core in association with The Stem Cell Institute will offer NGS on two platforms.
- ➢ HiSeq2000
  - > The HiSeq2000 can produce ~200billion bases (Gb) of sequence per run
- > GAIIx
  - > The GAIIx can produce up to 95billion bases (Gb) of sequence per run
- NGS Assays include:
  - RNA-Seq
  - ChIP-Seq
  - Exon capture and Sequencing
  - Targeted Resequencing
  - Whole Genome Sequencing
  - Microbiome
  - Methyl-Seq



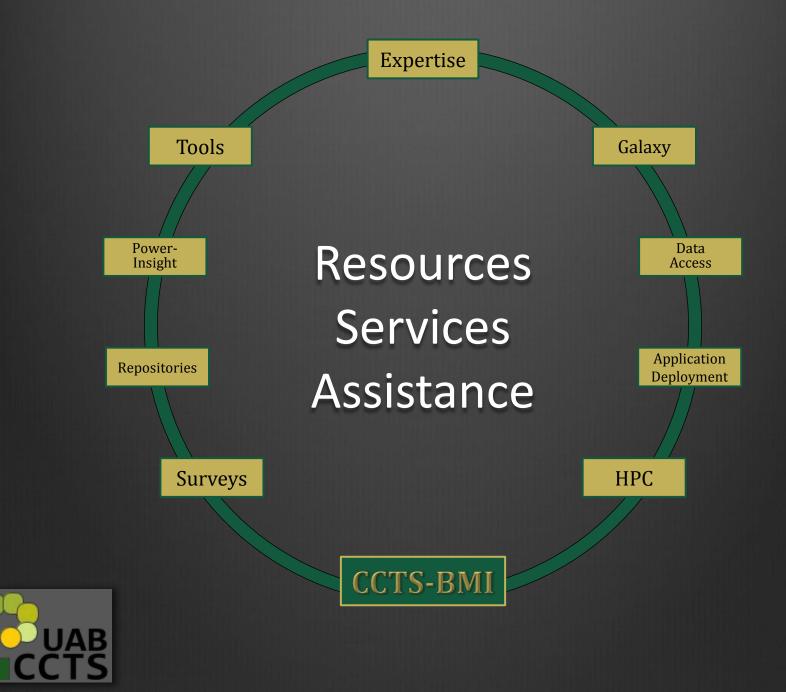


Cluster Generation <4 h (<10 min hands-on)





RTA v1.7, CASAVA v1.7 2 days (30 min hands-on)



# Genomics Pipeline at UAB

Study Design and Consultation

Sample Acquisition

- Laboratories
- Clinics
- Hospitals

Sample Processing

Logging and Storage
DNA/RNA extraction
Assay Preparation

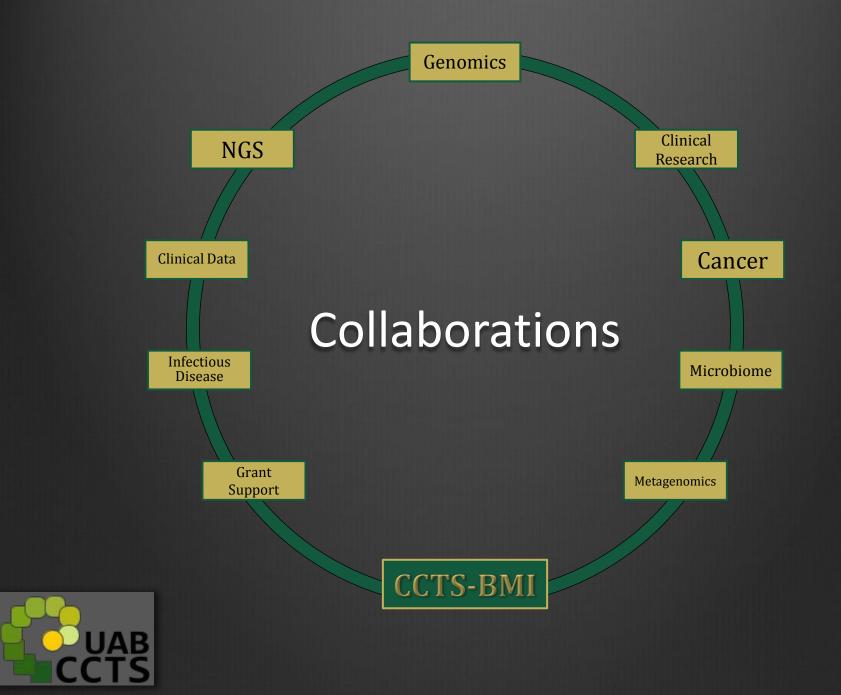
HEFLIN CENTER FOR GENOMIC SCIENCES Low to High Throughput Genomic and Genetic Assays; Focused Next Gen Sequencing and GX

> HUDSON ALPHA INSTITUTE Large Scale Next Gen and Specialized Sequencing Projects; High Volume Genomics Assays

#### Data Analysis

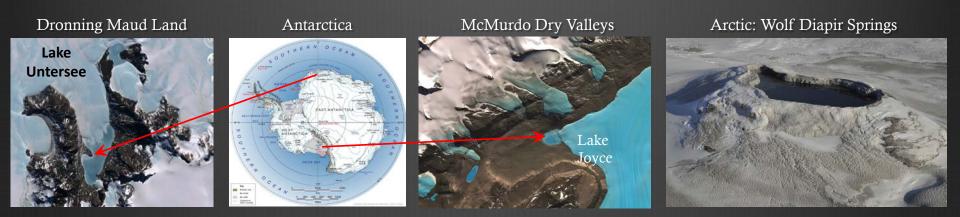
- Sequence analysis
- Gene expression analysis
- Genotyping analysis
- Methylation analysis

Knowledge Papers Grants



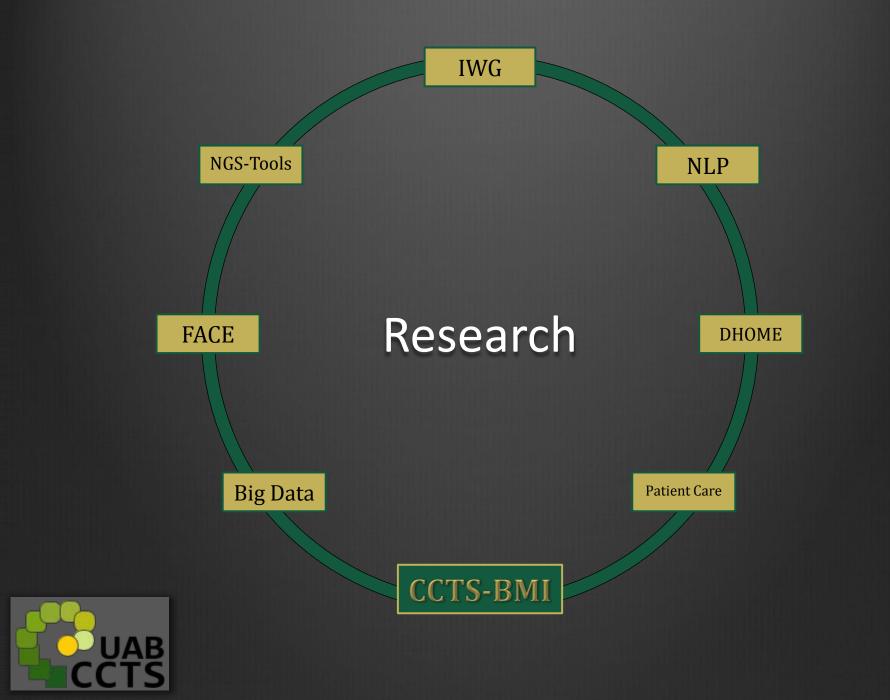
### Microbes in Extreme Cold Ecosystems: Diversity, Ecosystem Functioning and Bioprospecting

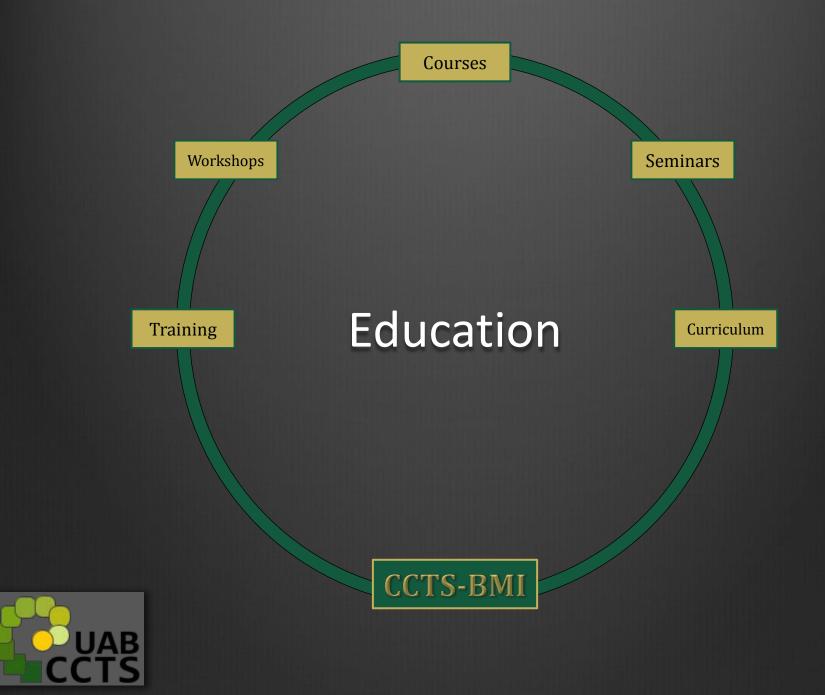
Jonathan Huang and Asim Bej, UAB Department of Biology; Ranjit Kumar, Elliot Lefkowitz, UAB Department of Microbiology; Mike Crowley, UAB Heflin Genomics Center; Dale Andersen, SETI Institute, CA



#### **OBJECTIVES:**

- **Ilumina sequencing:** Analyze genomic potential of the identity and function of microbes in Arctic and Antarctic perennially ice-covered freshwater lakes.
- **Comparative Metagenomics:** Deciphering potential biogeochemical cycling (*N*, *S*, *P*, *C*, *Fe*) and ecosystem functioning (*metabolic genes*) within and between these lakes.
- **Bioprospecting:** Mining enzymes and secondary metabolites for biomedical and pharmaceutical products.





### UAB Biomedical Informatics Resources and Assistance

- ➢ CCTS-BMI
  - Elliot Lefkowitz, Ph.D.
    - ElliotL@uab.edu
    - > (205) 934-1946
- Heflin Center
  - David Crossman, Ph.D.
    - ➢ dkcrossm@uab.edu
    - > (205) 996-4045

#### Section on Statistical Genetics (School of Public Health)

- Hemant Tiwari, Ph.D.
  - Htiwari@soph.uab.edu.
  - (205) 934-4907
- Department of Pathology Division of Informatics
  - Jonas Almeida, Ph.D.
    - jalmeida@uab.edu
    - (205) 975-3286
- Comprehensive Cancer Center (CCC) Biostatistics and Bioinformatics
  - Karan Singh, Ph.D.
    - kpsingh@uab.edu
    - > (205) 996-6122
- Microbiome Core Facility
  - Casey Morrow, Ph.D.
    - ▹ caseym@uab.edu
    - ► (205) 934-5705

